

CRF Errors Corrected by the STIC System Branch

Serial Number: 08/985474

CRF Processing Date: 11/10/98
 Edited by: 11
 Verified by: 11 (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:

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APR 21 2000

TECH CENTER 1600/2900

- ☒ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☒ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/12/98
TIME: 18:29:19

INPUT SET: S29791.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Klatzmann, David
6 Cohen, Jacques
7
8 (ii) TITLE OF INVENTION: a-B C4BP-TYPE RECOMBINANT
9 HETEROMULTIMERIC PROTEINS
10
11 (iii) NUMBER OF SEQUENCES: 6
12
13 (iv) CORRESPONDENCE ADDRESS:
14 (A) ADDRESSEE: SPENCER & FRANK
15 (B) STREET: 1100 New York Avenue, N.W., Suite 300 East
16 (C) CITY: Washington
17 (D) STATE: DC
18 (E) COUNTRY: USA
19 (F) ZIP: 20005
20
21 (v) COMPUTER READABLE FORM:
22 (A) MEDIUM TYPE: Floppy disk
23 (B) COMPUTER: IBM PC compatible
24 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
25 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
26
27 (vi) CURRENT APPLICATION DATA:
28 (A) APPLICATION NUMBER:
29 (B) FILING DATE:
30 (C) CLASSIFICATION:
31
32 (vii) PRIOR APPLICATION DATA:
33 (A) APPLICATION NUMBER: PCT/FR96/01132
34 (B) FILING DATE: 18-JUL-1996
35
36 (viii) ATTORNEY/AGENT INFORMATION:
37 (A) NAME: Gollin, Michael A.
38 (B) REGISTRATION NUMBER: 31,957
39 (C) REFERENCE/DOCKET NUMBER: GUPLA 0007
40
41 (ix) TELECOMMUNICATION INFORMATION:
42 (A) TELEPHONE: 202-414-4000
43 (B) TELEFAX: 202-414-4040
44
45
46 (2) INFORMATION FOR SEQ ID NO:1:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/12/98
TIME: 18:29:20

INPUT SET: S29791.raw

47
48 (i) SEQUENCE CHARACTERISTICS:
49 (A) LENGTH: 20 base pairs
50 (B) TYPE: nucleic acid
51 (C) STRANDEDNESS: single
52 (D) TOPOLOGY: linear
53
54 (ii) MOLECULE TYPE: other nucleic acid
55 (A) DESCRIPTION: /desc = "primer"
56
57
58
59
60 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
61
62 GAGACCCCCG AAGGCTGTGA 20
63
64 (2) INFORMATION FOR SEQ ID NO:2:
65
66 (i) SEQUENCE CHARACTERISTICS:
67 (A) LENGTH: 29 base pairs
68 (B) TYPE: nucleic acid
69 (C) STRANDEDNESS: single
70 (D) TOPOLOGY: linear
71
72 (ii) MOLECULE TYPE: other nucleic acid
73 (A) DESCRIPTION: /desc = "primer"
74
75
76
77
78 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
79
80 CTCGAGTTAT AGTTCTTTAT CCCAAGTGG 29
81
82 (2) INFORMATION FOR SEQ ID NO:3:
83
84 (i) SEQUENCE CHARACTERISTICS:
85 (A) LENGTH: 28 base pairs
86 (B) TYPE: nucleic acid
87 (C) STRANDEDNESS: single
88 (D) TOPOLOGY: linear
89
90 (ii) MOLECULE TYPE: other nucleic acid
91 (A) DESCRIPTION: /desc = "primer"
92
93
94
95
96 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
97
98 GCGGCCGCAG AGACCCCCGA AGGCTGTG 28
99

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/12/98
TIME: 18:29:21

INPUT SET: S29791.raw

100 (2) INFORMATION FOR SEQ ID NO:4:

101

102 (i) SEQUENCE CHARACTERISTICS:

103 (A) LENGTH: 23 base pairs

104 (B) TYPE: nucleic acid

105 (C) STRANDEDNESS: single

106 (D) TOPOLOGY: linear

107

108 (ii) MOLECULE TYPE: other nucleic acid

109 (A) DESCRIPTION: /desc = "primer"

110

111

112

113

114 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

115

116 CCACTTTGGA TAAAGAACTA TAA

23

117

118 (2) INFORMATION FOR SEQ ID NO:5:

119

120 (i) SEQUENCE CHARACTERISTICS:

121 (A) LENGTH: 36 base pairs

122 (B) TYPE: nucleic acid

123 (C) STRANDEDNESS: single

124 (D) TOPOLOGY: linear

125

126 (ii) MOLECULE TYPE: other nucleic acid

127 (A) DESCRIPTION: /desc = "primer"

128

129

130

131

132 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

133

134 ATTTCTAGAG AGTTATAGTT CTTTATCCAA AGTGGA

36

135

136 (2) INFORMATION FOR SEQ ID NO:6:

137

138 (i) SEQUENCE CHARACTERISTICS:

139 (A) LENGTH: 46 base pairs

140 (B) TYPE: nucleic acid

141 (C) STRANDEDNESS: single

142 (D) TOPOLOGY: linear

143

144 (ii) MOLECULE TYPE: other nucleic acid

145 (A) DESCRIPTION: /desc = "primer"

146

147

148

149

150 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

151

152 CCGGGACAGG TCCTGCTGGA ATCCAACATC AAGGTTCTGC CCACAG

46

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PAGE: 4

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474

DATE: 11/12/98
TIME: 18:29:22

INPUT SET: S29791.raw

153

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/983,474

DATE: 11/12/98
TIME: 18:29:22

INPUT SET: S29791.raw

Line

Error

Original Text

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/10/98
TIME: 16:35:25

INPUT SET: S29791.raw

N/A

This Raw Listing contains the General
Information Section and up to the first 5 pages.

Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Klatzmann, David
Cohen, Jacques(ii) TITLE OF INVENTION: a-B C4BP-TYPE RECOMBINANT
HETEROMULTIMERIC PROTEINS

(iii) NUMBER OF SEQUENCES: 6

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: SPENCER & FRANK
(B) STREET: 1100 New York Avenue, N.W., Suite 300 East
(C) CITY: Washington
(D) STATE: DC
(E) COUNTRY: USA
(F) ZIP: 20005

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: PCT/FR96/01132
(B) FILING DATE: 18-JUL-1996
(C) CLASSIFICATION:*I (vii) PRIOR APP DATA:*

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Gollin, Michael A.
(B) REGISTRATION NUMBER: 31,957
(C) REFERENCE/DOCKET NUMBER: GUPLA 0007

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 202-414-4000
(B) TELEFAX: 202-414-4040

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/10/98
TIME: 16:35:26

INPUT SET: S29791.raw

47 (C) STRANDEDNESS: single
48 (D) TOPOLOGY: linear
49
50 (ii) MOLECULE TYPE: other nucleic acid
51 (A) DESCRIPTION: /desc = "primer"
52
53
54
55

56 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
57

58 GAGACCCCCG AAGGCTGTGA
59

20

60 (2) INFORMATION FOR SEQ ID NO:2:
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62 (i) SEQUENCE CHARACTERISTICS:
63 (A) LENGTH: 29 base pairs
64 (B) TYPE: nucleic acid
65 (C) STRANDEDNESS: single
66 (D) TOPOLOGY: linear
67

68 (ii) MOLECULE TYPE: other nucleic acid
69 (A) DESCRIPTION: /desc = "primer"
70
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72
73

74 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
75

76 CTCGAGTTAT AGTTCTTTAT CCCAAGTGG
77

29

78 (2) INFORMATION FOR SEQ ID NO:3:
79

80 (i) SEQUENCE CHARACTERISTICS:
81 (A) LENGTH: 28 base pairs
82 (B) TYPE: nucleic acid
83 (C) STRANDEDNESS: single
84 (D) TOPOLOGY: linear
85

86 (ii) MOLECULE TYPE: other nucleic acid
87 (A) DESCRIPTION: /desc = "primer"
88
89
90
91

92 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
93

94 GCGGCCGCAG AGACCCCCGA AGGCTGTG
95

28

96 (2) INFORMATION FOR SEQ ID NO:4:
97

98 (i) SEQUENCE CHARACTERISTICS:
99 (A) LENGTH: 23 base pairs

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/983,474DATE: 11/10/98
TIME: 16:35:27

INPUT SET: S29791.raw

100 (B) TYPE: nucleic acid
101 (C) STRANDEDNESS: single
102 (D) TOPOLOGY: linear
103
104 (ii) MOLECULE TYPE: other nucleic acid
105 (A) DESCRIPTION: /desc = "primer"
106
107
108
109

110 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
111

112 CCACTTTGGA TAAAGAACTA TAA
113

23

114 (2) INFORMATION FOR SEQ ID NO:5:
115

116 (i) SEQUENCE CHARACTERISTICS:
117 (A) LENGTH: 36 base pairs
118 (B) TYPE: nucleic acid
119 (C) STRANDEDNESS: single
120 (D) TOPOLOGY: linear
121

122 (ii) MOLECULE TYPE: other nucleic acid
123 (A) DESCRIPTION: /desc = "primer"
124
125
126
127

128 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
129

130 ATTTCTAGAG AGTTATAGTT CTTTATCCAA AGTGGA
131

36

132 (2) INFORMATION FOR SEQ ID NO:6:
133

134 (i) SEQUENCE CHARACTERISTICS:
135 (A) LENGTH: 46 base pairs
136 (B) TYPE: nucleic acid
137 (C) STRANDEDNESS: single
138 (D) TOPOLOGY: linear
139

140 (ii) MOLECULE TYPE: other nucleic acid
141 (A) DESCRIPTION: /desc = "primer"
142
143
144
145

146 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
147

148 CCGGGACAGG TCCTGCTGGA ATCCAACATC AAGGTTCTGC CCACAG
149

46

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/983,474

DATE: 11/10/98
TIME: 16:35:28

INPUT SET: S29791.raw

Line	Error	Original Text
28	Wrong application Serial Number	(A) APPLICATION NUMBER: PCT/FR96/01132